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TGT	GTA	CAA	ATC	ATG	GGT	CGT	GGG	AAG	ATT	GAG	ATA	AAG	AGG	ATT	GAA	AAT	ACT	54
M	G	R	G	K	I	E	I	K	R	I	E	N	T					
ACG	AAC	CGA	CAG	GTC	ACT	TTC	TGC	AAG	CGC	CGA	AAT	GGT	TTA	TTA	AAG	AAG	GCG	108
T	N	R	Q	V	T	F	C	K	R	R	N	G	L	L	K	K	A	
TAT	GAA	TTA	TCA	GTT	CTT	TGT	GAT	GCA	GAA	GTG	GCC	CTC	ATC	GTC	TTC	TCC	AGC	162
Y	E	L	S	V	L	C	D	A	E	V	A	L	I	V	F	S	S	
AGA	GGG	AGA	CTT	TAT	GAA	TTT	GCC	AAC	CAC	AGC	GTG	AAG	AGG	ACG	ATT	GAG	AGG	216
R	G	R	L	Y	E	F	A	N	H	S	V	K	R	T	I	E	R	
TAC	AAG	AAG	ACT	TGC	GTT	GAC	AAC	CAC	GGA	GGG	GCG	ATA	TCA	GAG	TCC	AAT	270	
Y	K	K	T	C	V	D	N	N	H	G	G	A	I	S	E	S	N	
TCT	CAG	TAT	TGG	CAA	CAG	GAG	GCT	GGT	AAA	CTC	AGA	CAA	CAG	ATT	GAC	ATT	TTG	324
S	Q	Y	W	Q	Q	E	A	G	K	L	R	Q	Q	I	D	I	L	
CAA	AAT	GCA	AAT	AGG	CAT	TTG	ATG	GGT	GAC	GGG	CTT	ACA	GCT	TTG	AAC	ATT	AAG	378
Q	N	A	N	R	H	L	M	G	D	G	L	T	A	L	N	I	K	
GAA	CTC	AAG	CAA	CTT	GAG	GTT	CGA	CTT	GAA	AAA	GGA	ATC	AGC	CGA	GTG	CGA	TCC	432
E	L	K	Q	L	E	V	R	L	E	K	G	I	S	R	V	R	S	
AAA	AAG	AAC	GAG	ATG	TTG	CTT	GAA	GAG	ATC	GAC	ATC	ATG	CAG	AGA	AGG	GAA	CAC	486
K	K	N	E	M	L	L	E	E	I	D	I	M	Q	R	R	E	H	
ATA	CTT	ATC	CAG	GAG	AAT	GAG	ATT	CTT	CGC	AGC	AAG	ATA	GCC	GAG	TGT	CAG	AAT	540
I	L	I	Q	E	N	E	I	L	R	S	K	I	A	E	C	Q	N	
AGC	CAC	AAC	ACG	AAC	ATG	TTA	TCA	GCT	CCG	GAA	TAT	GAT	GCA	CTG	CCC	GCA	TTC	594
S	H	N	T	N	M	L	S	A	P	E	Y	D	A	L	P	A	F	
GAC	TCT	CGA	AAT	TTC	CTA	CAT	GCA	AAT	CTA	ATC	GAT	GCG	GCC	CAT	CAC	TAT	GCA	648
D	S	R	N	F	L	H	A	N	L	I	D	A	A	H	H	Y	A	
CAT	CAG	GAA	CAA	ACA	ACG	CTT	CAG	CTT	GGC	TGA	ACG	TTG	AAG	CGG	TGG	ACG	CTT	702
H	Q	E	Q	T	T	L	Q	L	G	*								
AAA	ACT	CAA	TCA	AGG	CAC	CCG	AAA	AAT	ATG	CTA	GTA	ACC	TTG	AAT	GAG	ATT	CAG	756
AGT	CGA	AAT	ATT	GCG	AGG	CAA	GAG	CAC	AAT	GGA	AGA	GAT	AGC	TCC	TAG	TAT	GAA	810
TAT	GGA	TTT	ATG	ATA	TTA	ACA	TAT	GGT	TTG	TCA	GCT	TTA	AAT	ATA	GCT	TTT	TGA	864
AAC	AAA	GAA	TAC	AAC	ATA	TTA	GCT	AGT	ATT	TTT	TTG	GCG	CAT	GTG	ATC	TTT	CTG	918
TTG																	921	

FIGURE 1

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AAACTCGACAGCAAATATGATTTAGATTATGACCTAGAAATAAGCATAGCATTAAAGCAT 60
ATACATAACAAGCGGTGATATACTCTGACTGCCACTGTACTTGAGGAAAGGTAGTGGACT 120
CTGCTCAGGTACATTAGTTGTAAGGTTGGCTTGGCTCTGGTAATATGAGAAAGTAAA 180
GAAGTAAAAGGTATTTGACTCTAGTCAAGTACATTGGATTGCCTTGTGGGGCTTGGAT 240
GGCTTGGGTTCGTGTGAGAAGCCAACAATTATAAGAAATATATAAAAATAAAAATAAAA 300
AAATTAAAGTGTGGAAGTGAAAACGGTGGGCAGAAATATACACAGAAGAGTACTTTAA 360
CAATGCGCAACCAAGGCAGATTACAACATTGATTCTGGACCTCGAATACGAGATAATGG 420
TGGTAAGAAATAAGGAAGAGTGGAGTGCATTGAAAATGAATGGAGAGCGCACAAAATG 480
GAGGACGAATAAAATGAAATATAATGCAAGAGTGCAATTCCCTATTATTCAGAAATGTA 540
TATGTGGGTCGGCATTACATGGCGTCGCATTCAAGGGGTGTCATAGCGGTCTTGA 600
TTGCAGTGTGGAGTTGCAACATGTACCAACAAATCCATTCACTCCAAAACCTAAATTAA 660
TCCTCTCCATTACTATTACCTACACCTACACCTAGTAAATATGTCCTGCCTTGTAACTCC 720
TCCACTGCCTGCACACGTCTTAGTCAATCCATCTGCCTCAAATAGGCATTATTTGTC 780
TTTCCCTCCGACTGAAAGGCTATCGACCGACCGACCGCTCATCTTCTTCTGCGCAA 840
TTTTTCTGCTGGATCATCATCATTACCATCATGCCATCCCCACCATCATCATGAT 900
GGTATCTCTATCTCTCCCTGGCAATCGATTGTAGAGGAAAGGAAGAGGGAAAGGGCATAT 960
GTATTGATCAACCTACCCGAAAAAAACAATCTGATCAGCCCTGCTCAATCTGCTTATAAA 1020
TCTCTTATCCACTGTTCAATCATTCAAGGTTCTCCACTTCAAGCAAAGGCGCCCGGA 1080
TTGGCCGTGTTCTAGATTCAGGTACTTAAATGGACAATATTCCCCACCTGAAGCCGT 1140
TCTGAAAAAGATTGTTGAGAAACAAACGATTGTAATATTGCTTAAGTTGAGCTTAA 1200
GGGGTTGGTACCTAACTTGCTTGTGGTTATTGTTCTCAGAAACTCGGCTGCGTCCA 1260
ACTGTAGGAACGAACCAGCACAGGGTTGCAGCTTGCTGTTGCGCCATTG 1320
CTTTTGGACTGGTATTAGTAGTTGCAGCTTGTGTCATACGCTGTGAGGATCTGTGCG 1380
CGGAAATTTGTGTACAAATC 1401

FIGURE 2

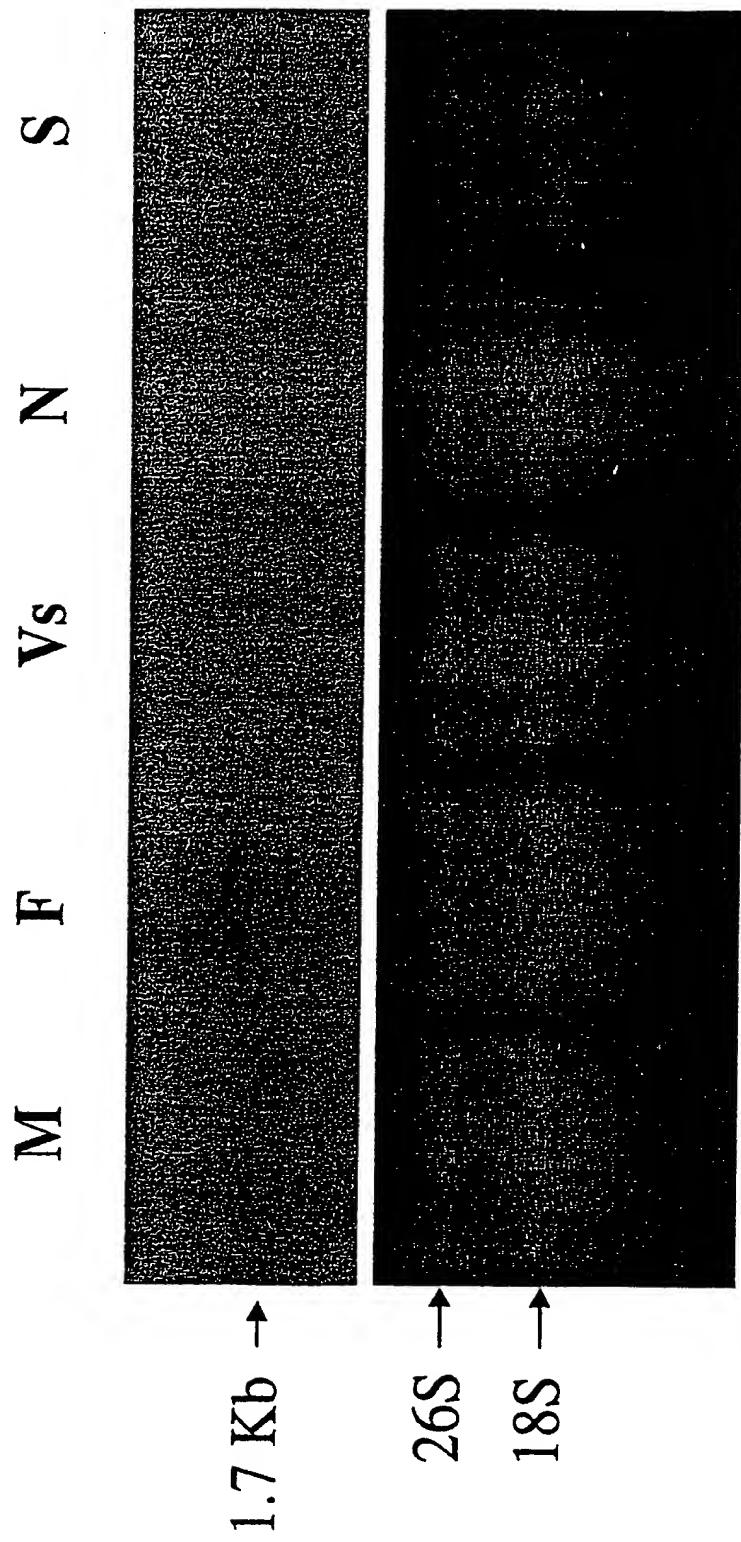
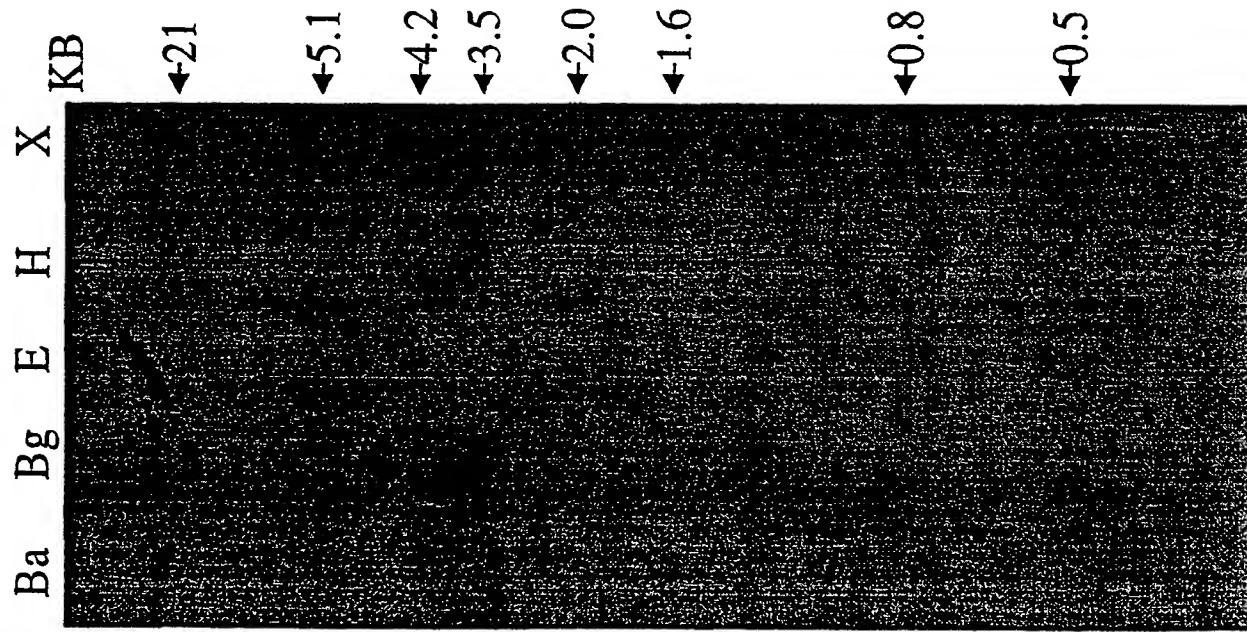


FIGURE 3

FIGURE 4



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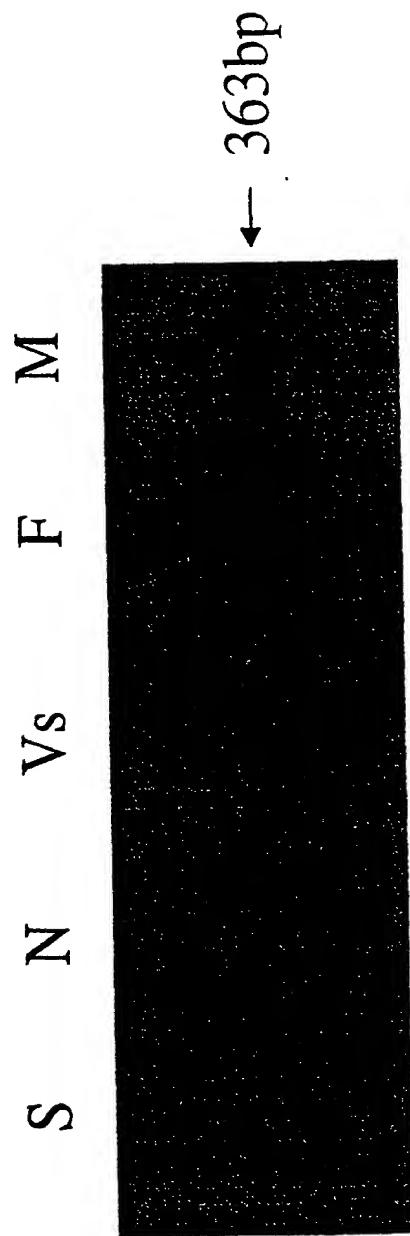


FIGURE 5

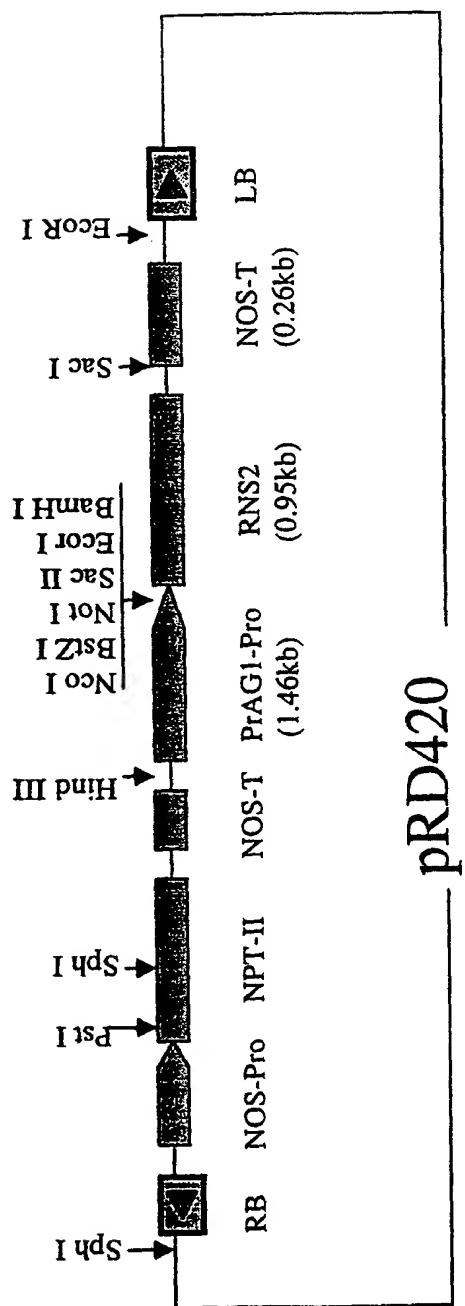


FIGURE 6

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Figure 7

PC C1 C2 C3 1a 1b 2 3a 3b 3c 4a 4b

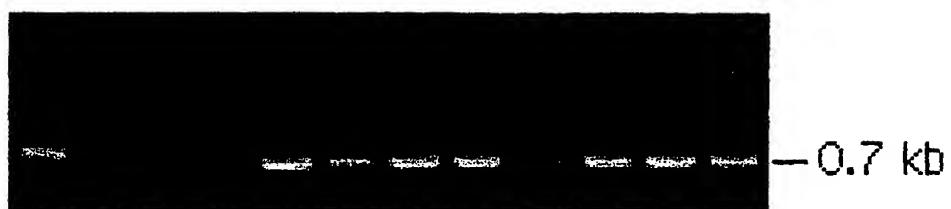


Figure 8

C1 C2 C3 1a 1b 2 3a 3b 3c 4a 4b

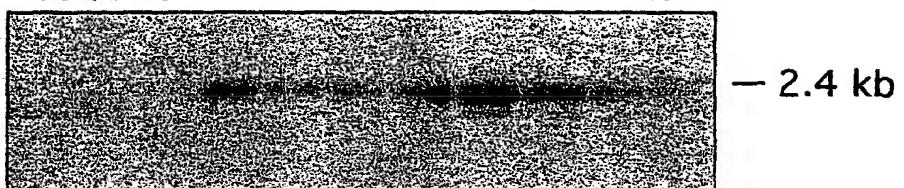


Figure 9

C1 C2 C3 1a 1b 2 3a 3b 3c 4a 4b

